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Q96dk0
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Q9d814
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
NON_TER 124
SEQUENCE 124 AA; 13580 MW;
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Q96QS0;
01-DEC-2001
01-DEC-2001
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
Young D.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                       Clin. Immunol. Immunopathol. 87:184-192(1998). EMBL; AF035022; AAD56258.1; -. HSSP; P01751; INQB.
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111 AVSS 114 82; Conservative h 69.9%; Score 416.5; DB 2; Length 124; Similarity 66.1%; Pred. No. 4.2e-36; B2; Conservative 12; Mismatches 19; Indels 11 AQKFQGRVIMIRDISTSTVYMELSSLRSEDTAVYYCARGLYVVVPAAFSRFDYWGQGTLV APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC------LLQALKHWGQGTLV 110 (TrEMBLrel. 19, Created)
(TrEMBLrel. 19, Last sequence update) PRELIMINARY; 13580 MW; 1BAAACBD96ACD2A2 CRC64; PRT; 159 ₿ 11; Gaps 60

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RESULT 3
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             GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                     EMBL; J00240; AAA52988.1; -. PIR; A02024; HVHUHG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=83144028; PubMed=6298778;
Rechavi G., Ram D., Glazer L., Zakut R., Givol
"Evolutionary aspects of immunoglobulin heavy."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Proc. Natl. Acad. Sci. U.S.A. 80:855-859(1983).
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(Rel. 42, Last ann
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; IG LIKE; 1.
A; 17497 MW;
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60.0%;
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annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 390.5; DB 2;
Pred. No. 3.1e-33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL/GenBank/DDBJ
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           precursor.
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Best Local S
Matches 71
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                                                                                                                                                                              InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; Ig_LIKE; 1.
NON TER
NON_TER 125 125
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CHAIN 20
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Q9UL95;
01-MAY-2000
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NON TER
SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Immunoglobulin V region; Sign
                                                                                                                                                                                                                                                                                                                                                                                               Myosin-reactive (Fragment).
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01-OCT-2003
                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                              EMBL; AF035019; AAI
HSSP; P01751; 1NQB
                                                                                                                                                                                                                                                                              fetus.
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                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                           WEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S
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121
                    110 VAVSS
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                                                                                             1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                  N
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VIVSS
                                         AQKVQGRVTMTRDTTISTAYMELSRLRSDDTAVYYCARSQGGGRIAAAGDAFDIWGQGTM
                                                             APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLL------
                                                                                VQLVE-SGAEVKKPGASVKVSCKASGYTFTGYYMHWVRQAPGQGLEWMGWINPNSGGTNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AQKFQGRVIMTRDTSTSTVYMELSSLRSEDTAVYYC 115
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117
117
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125 AA;
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
rive immunoglobulin heavy chain variable
                                                                                                                            Conservative
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                    114
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117
                                                                                                                                                                      13516 MW;
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                                                                                                                                      62.4%;
58.4%;
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                                                                                                                            15;
                                                                                                                                      Score 372; DB 2;
Pred. No. 2.1e-31;
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Ig-like.
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                                                                                                                            Mismatches
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                                                                                                                                               Length 125;
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3.M.,
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RESULT 5
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ID Q6N095

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Best Local !
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SMART; SM00407; IGC1; 3.
SMART; SM00406; IGV1;
SMORT; SM00406; IGV 1.
PROSITE; PS00835; IG LIKE; 4
PROSITE; PS00890; IG_MHC; UN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig-cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; Cl-set; 3.
                                                        (1)
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934; I
MEDLINE=98277139; Van der Merwe P.L.,
                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                             Young
"Myos
                                                                                                                                                                                  01-MAY-2000
01-OCT-2003
                                                                                                                                                                                                         Q9UL94;
01-MAY-2000
                                                                                                      NCBI_TaxID=9606;
[1]
                                                                                                                                                                      Myosin-reactive
                                                                                                                                                                                                                                 Q9UL94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical SEQUENCE 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The German Human cDNA Consortium; Wambutt R., Heubner D., Mewes H.W., Weil B., Amid C., O: Fobo G., Han M., Wiemann S.; Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases EMBL; BX640621; CAE45775.1; -. HSSP; P01861; 1ADQ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686K03196.
Name=DKFZp686K03196;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
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Immunol. Immunopathol. AF035020; AAD56256.1; P01751; 1NQB.
                                             in-reactive
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77; Conserv
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nilarity 61.1%;
Conservative 10
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
tive immunoglobulin heavy chain variable
                                                                                                                                                                                                                               PRELIMINARY;
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                                             autoantibodies
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; UNKONOWN_2.
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                      87:184-192(1998)
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Pred. No. 1.8e-30;
D; Mismatches 26;
                                                                                                                        Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Catarrhini; Hominidae;
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                                                                DOI=10.1006/clin.1998.
, Kalis N.N., Berney S.
                                            rheumatic
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                                             and normal
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Best Local
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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl:
InterPro; IPR003597; Ig_MC.
InterPro; IPR003596; Ig_W.
Pfam; PF07654; Cl-Bet; 2.
SMART; SM00409; IG; 4.
SMART; SM00407; IGC1; 3.
SMART; SM00406; IGV; I.
PROSITE; PS00835; IG_LITE; 4.
PROSITE; PS00835; IG_MHC; UNKW.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Human rectum tumor;
The German Human cDNA Consortium;
The German Human cDNA Consortium;
Poustka A., Albert R., Moosmayer P., Schupp I., Wellenreuther R.,
Mewes H.W., Weil B., Amid C., Osanger A., Fobo G., Han M., Wiemann
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BX640710; CAE45829.1; -.
HSSP; P01751; 1A6W.
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NON_TER
SEQUENCE
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NON_TER 1
SEQUENCE 498 AA; 5
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Mammalia; Eutheria;
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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155
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                                                                                                                                                                                                      VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
TVSS
                                                                                                                            APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQA-----LKHWGQGTLV
                                                                                                                                                                         VQLV-QSGADVKKPGASVKVSCKASGYTFTNYFFHWVRQAPGQGPEWMGMINPRDGSTKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQA-----LKHWGQGTLVAVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                    AQRFQGRVSMTRDTSTSTIYMELSSLRSEDTAMFFCARAGPGYGTSASYYFDYWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AQKFQGKVTMTKDTSISTAYMELSRLRSDDTAVYYCARGGGRGLWFDPWGQGTLVTVSS
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119 AA;
                                                                                                                                                                                                                                                                   Conservative
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13205 MW;
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Primates;
                                                                                                                                                                                                                                                                                                                                                            54125 MW;
                                                                                                                                                                                                                                                                                        61.2%;
56.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 UNKNOWN
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Pred. No. 4.2e
16; Mismatches
                                                                                                                                                                                                                                                                Score 364.5; DB 2;
Pred. No. 6.5e-30;
0; Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vo
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                            40B3208A84E03B46 CRC64;
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i; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2;
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on update)
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HV1C_HUMAN
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Best Local (
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Bell L.O., Gould H.J.;
"Cloning and sequence determination of the gene rimmunoglobulin epsilon chain expressed in a myeldimunoglobulin expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOMAIN
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SEQUENCE FROM N.A.
MEDLINE=83065234; PubMed=6815656;
MEDL-- " H Molgaard H.V., Houghton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                901744;
21-JUL-1986 (Rel. 01, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation updat
Ig heavy chain V-I region ND precursor (Fra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSSP, P01751; NOB.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR0037110; Ig-like.
InterPro; IPR003756; Ig V.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennich H.H., Johansson S.G.O., von Bahr-Lin (In) Bach M.K. (eds.); Immediate hypersensitivity: modern concepts Marcel Dekker, New York (1978).

-I- MISCELLANEOUS: This epsilon chain was is
      Q65ZC8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS50835; IG_LIKE; 1.
Direct protein sequencing; Immunoglobulin
Pyrrolidone carboxylic acid; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pyrrolidone
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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SIMILARITY:
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                                                                                                                                                                                                          VAVSS 114
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                                                                                                                                                   VIVSS
                                                                                                                                                                                                                                                                     FOGRVTMTRDASFSTAYMDLRSLRSDDSAVFYCAKSDPFWSDYYNFDYSYTLDVWGQGTT
                                                                                                                                                                                                                                                                                                                         FKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQ---
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147 AA;
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   PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15;
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IH -> HI (in Ref. 2).
VG -> GV (in Ref. 2).
Missing (in Ref. 2).
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Ig-like.
Pyrrolidone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 356;
Pred. No. 1.
      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             948F9F72A5366C20 CRC64;
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r (Fragments).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 147;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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cell line.";
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                                                                                                                                                                                                                                                                                                                               -ALKHWGQGTL
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RESULT 10
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Best Local S
Matches 68
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InterPro; IPR007110; Ig-like.
InterPro; IPR003506; Ig_v.
Pfam; PF00047; ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q6N030;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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25-OCT-2004 (7
25-OCT-2004 (7
25-OCT-2004 (7
Single-chain |
                                 InterPro;
InterPro;
                                                                                                                  InterPro;
InterPro;
                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                 The German Human cDNA Consortium;
Poustka A., Albert R., Moosmayer P., Schupp I., Wei
Mewes H.W., Weil B., Amid C., Osanger A., Fobo G.,
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ datal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein DKFZp686I15212.
Name=DKFZp686I15212;
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Mammalia; Eutheria;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q6N030
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kontermann R.E., Wing M.G., Winter G.; "Complement recruitment using bispecific Nat. Biotechnol. 15:629-631(1997). EMBL; Y13057; CAA73500.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=97362799;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=scFv;
                                                                                                                                                                                                                                                                                                                                                         TISSUE=Human rectum tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                          InterPro;
                                                                                    ; BX640724; CAE45841.1; -.

7; P01861; 1ADQ.

rrPro; IPR000005; HTHAraC.

rrPro; IPR007310; Ig-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AQRFQGRVTMTRDTSISAAYMEVSRLRSDDTAVYYCAREGTGSAIYGMDVWGQGTLVTVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC-----LLQALKHWGQGTLVAVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VQLV-QSGAEVKKPGDSVKVSCKASGYTFSDHYMHWVRQAPGQGLEWMGWIDPNNGDTRF
                                    IPR003597;
IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
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(TrEMBLrel.
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n Fv (Fragmen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
Primates;
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Ig.
Ig-like.
Ig_cl.
Ig_MHC.
Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence up
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Ve
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         352; DB 2
No. 6e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vertebrata; Euteleostomi; i; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB
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RESULT 12
Q8WY24
ID Q8WY2
AC Q8WY2
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
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Best Local S
Matches 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 68
                                                                                                                               Q8WY24;
Q8WY24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NON_TER
NON_TER
SEQUENCE
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sMART; SM00409; IGc1; 3.
sMART; SM00407; IGC1; 3.
SMART; SM00406; IGV; 1.
PROSITE; PS00041; HTH ARAC FAMILY 1; UNKNOWN_1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_MHC; UNKNOWN_2.
  Homo sapiens (Human)
Eukaryota; Metazoa;
                                         SNC66 protein
                                                                    01-MAR-2004
                                                                                     01-MAR-2002
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. Song X.T., Feng Z.Q., Guan Y Submitted (JUN-2000) to the EMBL; AF282622; AAG01452.1; HSSP; P01751; 1A6W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schistosoma japonicum (Blood fluke).
Eukaryota; Metazoa; Platyhelminthes;
Schistosomatoidea; Schistosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
01-OCT-2003 (TrEMBLrel. 25,
Monoclonal anti-idiotypic ar
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01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=6182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9GYZ2
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SEQUENCE 5
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                                                                                                                                                                                                                                                                            APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLL-----QALKHWGQGTLVAVSS
                                                                                                                                                                                                                                                                                                                                                                  VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                                                                                                                                                        NQKFKDRVTMTTDKSFSTAYMDLRSLRSADSAVYYCARYYDDHYCLDYWGQGTTVTVSS
                                                                                                                                                                                                                                                                                                                                     VQLVE-SGAEVRKPGASVRVSCKASGYTFTGYYMNWVRQAPGHGLEWIGYINPSRGYTNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PS50835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQA-----LKHWGQGTLVAVSS 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYAPK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            119
119 AA;
                                                               2 (TrEMBLrel.
2 (TrEMBLrel.
4 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       al protein.
518 AA; 5
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nilarity 57.1%;
Conservative 1:
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                                                                                                                                                   PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            119
13567 MW;
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Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                            58.2%;
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5, Last annotation update)
antibody NP30 heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                             13;
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e EMBL/GenBank/DDBJ
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                                                           Created)
Last sequence that the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 347; DB 2;
Pred. No. 8.9e-29;
.3; Mismatches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 348; DB
Pred. No. 3.8e
L5; Mismatches
  Craniata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BA893873FD5FA6AB
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                                                             sequence up
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Schistosoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.8e-28;
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                                                             on update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 518;
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  Euteleostomi;
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Q924Q1
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                                                                                                                                                                                             Query Match
Best Local S
Matches 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kozono Y., Kozono H., Azuma T
Submitted (AUG-2001) to the E
EMBL; AB069913; BAB63929.1; -
PIR; F33932; F33932
PIR; I28833; I28833.
PIR; PH0985; PH0985.
PIR; PH1155; PH1155.
PIR; PH1156; PH1156.
PIR; PH1157; PH1157.
PIR; PH1157; PH1158.
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SEQUENCE FROM N.A.

Zheng S., Shao X., Cao J., Geng L.,
Submitted (JUN-2000) to the EMBL/Ger

EMBL, AF283666; AAL36987.1; -.

HSSP; P01876; 10W0.

Pfam; PF07654; C1-set; 2.

SMART; SM00406; IGy; 1.

PROSITE; PS50835; IG LIXE; 4.

PROSITE; PS00290; IG MHC; UNKNOWN 1.

SEQUENCE 497 AA; 53665 MW; F24D0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q924Q1 PRELIMATION OF THE PRESENTING OF THE PROPERTY OF THE PR
                                                                                                                                                                                                                                                                                                                           PROSITE;
NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=C57BL/6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
[1]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P01751; 1A6W
64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                83
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                                                                                                                                                                                                                                                                                                                                                                                              SM00406; IGv; 1.
E; PS50835; IG_LIKE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23
                                                                                                                                                                                             Similarity
63; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4
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67; Conserv
                                                                                                  LEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYAPK
FKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCL---LQALKHWGQGTLVAVSS 114
                                                                LQQPGTELVKPGASVKLSCKASGYTFTSYWMHWVKQRPGQGLEWIGNINPSNGGTNYNEK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SS 144
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SS 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FOGRETFSROTSINTAYMVESSESTEDSALYFCARGNERGGRGFGYNWFDPWGHGTEVTV
                                                                                                                                                                                                                                                                                                                           142
142 AA;
                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                     55.3%;
                                                                                                                                                                                                                                                                                                                              15622 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54.9%;
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                                                                                                                                                                                                 19;
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e EMBL/GenBank/DDBJ
                                                                                                                                                                                          Score 340.5;
Pred. No. 5.3e
19; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T.;
EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 341.5; DB 2;
Pred. No. 1.8e-27;
7; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; 1
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                           24A265CE4EA4318B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F24D08DFA5A663E5
                                                                                                                                                                                          3; DB 2;
5.3e-28;
hes 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dong Q.;
J databases
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                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11;
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                                                                                                                                63
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RESULT
Q9UL89
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HV1G_HUMAN
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Best Local S
Matches 70
                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1991
01-NOV-1991
10-OCT-2003
                                       SEQUENCE FROM N.A.

MEDLINE=88296408; PubMed=2841108;

Matsuda F. Lee K.H., Nakai S., Sato T., Kodair

Matsuda F. Lee K.H., Nakai S., Sato T., Kodair

Ohno H., Fukuhara S., Honjo T.;

"Dispersed localization of D segments in the hu

heavy-chain locus.";

EMBO J. 7:1047-1051(1988).

EMBO J. 7:1047-1051(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wu X., L
  This
                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HV1G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; AF035025; AAD56261.1; PIR; PH0870; PH0870. PIR; PH1671; PH1671.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Myosin-reactive autoantibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS50835; IG_LIKE; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HSSP; P01751; 1NQB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Fragment)
                                                                                                                                                                                                                                                                                                                                                                            -NOV-1991 (Rel. 20, Created)
-NOV-1991 (Rel. 20, Last sequence update)
-OCT-2003 (Rel. 42, Last annotation update)
heavy chain V-I region V35 precursor.
SWISS-PROT entry is copyright.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYAPKFK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQ-----ALKHWGQGTLVAVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGRIIPILGIANYAQKFQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            116
116 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      57.0%;
llarity 60.9%;
Conservative 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 340; DB 2; I
Pred. No. 4.8e-28;
.0; Mismatches 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87:184-192(1998).
                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; l
Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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  19
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                                                                                                                                                                            Kodaira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽
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produced through a collaboration
                                                                                                                             human
                                                 domain.
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                                                                                                                                                                                                                                                                                                                               Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    region
                                                                                                                                                                          Zong
                                                                                                                                                                            8.0.,
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RESULT 2 S36265 S36265 Ig heavy chain V region (clone al C; Species: Homo sapiens (man) C; Dete: 03-Feb-1994 #sequence_rev C; Accession: S36265 R; Griffiths, A.D.; Malmqvist, M.; EMBO J. 12, 725-734, 1993 A; Title: Human anti-self antibodi A; Reference number: S36256; MUID: A; Accession: S36265 A; Status: preliminary; nucleic ac A; Molecule type: mRNA A; Residues: 1-118 <gri> A; Cross-references: EMBL: Z18846; C; Superfamily: immunoglobulin V r</gri>	chair: - Hom	P	344.5 57.8 132 344.5 57.8 132 344.5 57.8 132 344.5 57.7 108 343.57.6 135 343.57.6 627 342.5 57.3 136 341.5 57.3 136 341.5 57.3 136 341.5 57.1 132 340.5 57.1 132 340.5 57.1 132 340.5 57.1 132 340.5 57.0 117 339.5 57.0 117
pha-MUC1-1) - human (fision 03-Feb-1994 #tex Marks, J.D.; Bye, J.M Marks, J.D.; Bye, J.M 1918 specificies with high specificies 19178448; PMID:7679990 id sequence not shown id sequence provides 19178448; PIDN:CAA79 egion; immunoglobulin	(fragment) (fragment) (gragment) (gragm	ALIGNMENTS	PH0955 2 B33548 2 PH0954 2 PH1664 2 PH1664 2 PH0953 3 S1596 2 S26912 2 PH0960 2 S26912 2 PH0960 2 S26912 2 PH0960 2 S276912 2 PH0960 2 S276912 2 PH0960 3 S31596 3 PH1665 3 S31596 3 PH1665 3 S31596
(fragment) ext_change 23-Jul-1999 .M.; Embleton, M.J.; McCafferty, J city from phage display libraries. 90 n homology	tt_change 16-Aug-1996 uson, M.; Fry, K.E.; Borrebaeck, C uson, M.; Fry, M.; Borrebaeck, C uson, M.; Borrebaeck, C uson, M.; Borrebaeck, C		Ig heavy chain V r

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RESULT 4
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C;Date: 01-Feb-1995 #sequence_revision 12-May-1995 #text_change C;Accession: S49530
C;Accession: S49530
R;Mahmoudi, M.; Edwards, J.; Cairns, E.; Bell, D. submitted to the EMBL Data Library, October 1994
A;Description: Molecular characterization of natural human anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change C;Accession: $20783 C;Accession: $20783 C;Accession: $20783 C;Accession: $20783 C;Accession: $20764 A;Reference number: $20764 A;Accession: $20783
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                                                                    A;Cross-references: EMBL:Z46348; NID:g560839; PIDN:CAA86467.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology F;34-117/Domain: immunoglobulin homology <IMM>
                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-135 <MAH>
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C; Species: Homo sapiens (man)
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65.2%; bc.
64.7%; Pre
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62.0%; Pre
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     Score 388.5;
Pred. No. 3.4e
14; Mismatches
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Pred. No. 1.8e-29;
6; Mismatches 22
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Pred. No. 3.5
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A;Status: preliminary
A;Molecule type: DNA
A;Residus: 1-98 <-TOMS
A;Cross-references: EMBL:Z12309; NID:g32958; PIDN:CAA78179.1;
C;Superfamily: immunoglobulin V region; immunoglobulin homolog
C;Keywords: heterotetramer; immunoglobulin
E;15-98/Domain: immunoglobulin homology <-IMM>
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R;Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, J. Mol. Biol. 227, 776-798, 1992
J. Mol. Biol. 227, 776-798, 1992
A;Title: The repertoire of human germline V(H) sequences reveals aboa;Reference number: S26885; MUID:93021117; PMID:1404388
A;Accession: S26920
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A;Title: Developmentally restricted immunoglobulin heavy chain variable A;Reference number: A33548; MUID:89345575; PMID:2503826
A;Accession: D33548
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
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A; Residues: 1-123 <KIP>
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C;Date: 17-Jan-1990 #sequence_revision 17-Jan-1990 #text_change 16-Aug-1996
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                                         VQLV-QSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGIINPSGGSTSY
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Pred. No. 3.2e-28;
L2; Mismatches 12
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Pred. No. 6.5e-
.7; Mismatches
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F;34-117/Domain:
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A;Molecule type: mRNA
A;Residues: 1-126 < ZEB>
A;Cross-references: GB:M88309; NID:g183952; PIDN:AAA35967.1; PID:g183953
A;Note: nucleotide translation not given
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Title: Human combinatorial antibody libraries to hepatitis B surface antigen A;Reference number: A44151; MUID:92228746; PMID:1373487
A;Accession: I44151
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C;Date: 27-Jun-1994 #sequence_revision 27-Jun-1994 #text_change
C;Accession: I44151
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A;Title: Evolutionary aspects of immunoglobulin heavy chain variable A;Reference number: A02024; MUID:83144028; PMID:6298778
A;Accession: A02024
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A;Map position: 14q32.33-14q32.33
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A; Residues: 1-117 < REC>
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;Date: 03-Aug-1984 #sequence_revision 03-Aug-1984 #text_change 09-Jul-2004
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APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQ------ALKHWGQGTLVA 111
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                                                     VKLLE-SGAEVRKPGASVKVSCKASGYTFTGYYLHWVRQAPGQGLEWMGWIHPNNGGTFH
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                                                                                                                                                                                               immunoglobulin homology <IMM>
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                                                                                                                               Conservative
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                                                                                                                                            62.8%;
58.5%;
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                                                                                                                         18;
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                                                                                                                                            Score 374; DB 2
Pred. No. 7e-28;
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Pred. No. 3.8e-28;
2; Mismatches 12
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                                                                                                                                                          Length 126
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A; Residues: 1-137 < AND>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R; Andris, J.S.; Johnson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Accession:
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Best Local (
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Best Local :
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                                    106 QGTLVAVSS 114
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   KGTTVTVSS 129
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A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-129 <FIG>
A;Cross-references: EMBL:Z31680; NID:g509786; PIDN:CAA83485.1; PID:g1335146
A;Cross-references: EMBL:Z31680; viperiaminoglobulin homology
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Figini, M; Marks, J.D.; Winter, G.; Griffiths, A.D. J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains A;Reference number: S46390; MUID:94254092; PMID:8196048
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(41287)
Ig heavy chain precursor V-I region (71-31) (anti-HIV-1) - human
C;Species: Homo sapiens (man)
C;Date: 21-Apr-1992 #sequence_revision 21-Apr-1992 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: S46390;
A; Accession: S46393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Homo sapiens (man)
C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: GB:M67502; NID:g185409; PIDN:AAA52943.1; PID:g185410 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Reference number: A41287; MUID:91352074; PMID:1909030 A;Accession: C41287
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A; Title: Molecular characterization of five human anti-human
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61 AQKFQGWVTMTRDTSISTAYMELSRLRSDDTAVYYCARDSAYYYDSSGYYSANYYMDVWG
                         61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKH----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                   VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOKFOGRVAMTROTSTSAVYMELSSLRSVDTAIYYCAGVOGVARPLGDWGOGSLVTVSS 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VQLL-QSGAEVRKPGASVKISCKPSGFSFITYYFHWVRQAPGQGLEWMGVINPSGGT-TY
                                                                                          VQLV-QSGAEVKKPGASVKVSCKASGYTFTGYYMHWVRQAPGQGLEWMGWINPNSGGTNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPKFRGRVTMTRDTSISTAYMELSSLRSDDTAVYYCAIEYGDTDCGGDCYSIWGQGTLVT 121
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Pred. No. 3.8e
15; Mismatches
                                                                                                                                                                                      Score 365; DB 2;
Pred. No. 4.9e-27;
6; Mismatches 25
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C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 16-Aug-1996
C;Accession: PH1669
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 16-Aug-1996
C;Accession: PH1668
R;Hillson, J.L.; Karr, N.S.; Oppliger, I.R.; Mannik, M.; Sasso, E.H.'
J. Exp. Med. 178, 331-336, 1993
A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding A;Reference number: PH1642; MUID:93301610; PMID:8315388
A;Accession: PH1668
R;Olee, T.; Lu, E.W.; Huang, D.F.; Soto-Gil, R.W.; Deftos, M.; Kozin, J. Exp. Med. 175, 831-842, 1992
A;Title: Genetic analysis of self-associating immunoglobulin G rheuman
                                                                  Ig heavy chain V region precursor - hum (Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision C;Accession: $23623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: mRNA
A;Residues: 1-109 <HIL>
A;Experimental source: B cell
A;Experimently: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;7-90/Domain: immunoglobulin homology <IMM>
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PH1668
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A; Residues: 1-110 <HIL>
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Best Local :
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                                                                                                                                                                                                                                                     TMTRDTSISTAYMELSRLRSEDTAVYYCASLYGDYVDDWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                               AEVKKPGASVKVSCKASGYTFTGYYMHWVRQAPGQGLEWMGWINANSGGTNYAQKFQGRV
                                                                                                                                                                                                                                                                                                                                                                                  AEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYAPKFKGRL
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                                                                                                                                                                                                                                                                                               TMSRDSSTDTVYMTLTSLTSEDTAVYYCLL---QALKHWGQGTLVAVSS 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TMTRDTSISTAYMELSRLRSEDTAVYYCARDLTGDAFDIWGQGTMVTVSS
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self-associating immunoglobulin G rheumatoid factors
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 361.5;
Pred. No. 8.8e
12; Mismatches
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Pred. No. 8e-27;
                                                                                                                                      human
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                                                                                         13-Jan-1995
                                                                                                                                      (fragment)
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ches 25;
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                                                                                       #text_change 23-Jul-1999
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                                            F.; Carson,
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F;31-35/Region: complementarity-determining F;36-69/Region: framework 2 F;55-67/Region: complementarity-determining F;68-98/Region: framework 3 F;99-117/Region: complementarity-determining
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A;Molecule type: DNA
A;Residues: 1-129 <MARP>
C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Residues: I-129 <KIP>
R;Martin, T.; Duffy, S.F.; Carson, D.A.; Kipps, T.J.
J. Exp. Med. 175, 983-991, 1992
A;Title: Evidence for somatic selection of natural autoantibodies.
A;Reference number: PH0952; MUID:92202880; PMID:1552291
A;Accession: PH0956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Kipps, T.J.; Tomhave, E.; Pratt, L.F.; Duffy, S.; Chen, Proc. Natl. Acad. Sci. U.S.A. 86, 5913-5917, 1989
A;Title: Developmentally restricted immunoglobulin heavy A;Reference number: A33548; MUID:89345575; PMID:2503826
A;Accession: A33548
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A;Accession: S23623
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-171 <OLE>
A;Cross-references: EMBL:X59702; NID:g32010; PIDN:CAA42223.1; PID:g32011
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;34-117/Domain: immunoglobulin homology <IMM>
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C; Accession: A33548; PH0956
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Matches
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                                                                 AQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYCARGPRLLADVLLWFGELSEFDYWG
                                                                                         APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCL------
                                                                                                                                               VQLV-QSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGGIIPIFGTANY
                                                                                                                                                                                         VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
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Pred. No. 2.2e-26;
2; Mismatches 26
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121 QGTLVTVSS 129

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PH0982

Ig heavy chain V region (GG+ T-142) - human (fragment)

C; Species: Homo aspiens (man)

C; Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Aug-1996

C; Accession: PH0982

R; Martin, T; Duffy, S.F.; Carson, D.A.; Kipps, T.J.

J; Exp. Med. 175, 983-991, 1992

A; Title: Exidence for somatic selection of natural autoantibodies.

A; Reference number: PH0982; MUID:92202880; PMID:1552291

A; A; Carson, PH0982

A; Carson, PH0982; MUID:92202880; PMID:1552291

A; Carson in PH0982

A; Carson in PH0982; MUID:92202880; PMID:1552291

A; Carson in Ph0982; MUID:92202880; PMID:1552
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Maximum Match 100%
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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596
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Copyright (c) 1993 - 2005 Compugen Ltd
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US-08-477-989B-94
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US-08-234-180C-7
US-08-234-180C-7
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Query Match 66.5%;
Best Local Similarity 64.2%;
Matches 77; Conservative 1
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Result No.

14; Mismatches Score 396.5; DB 3 Pred. No. 1.8e-33;

Indels Length 120;

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Gaps

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363	363	364.5	364.5	364.5	364.5	364.5	364.5	364.5	364.5	366.5	366.5	367.5	367.5	368	368	368.5	368.5
60.9	60.9	61.2	61.2	61.2	61.2	61.2	61.2	61.2	61.2	61.5	61.5	61.7	61.7	61.7	61.7	61.8	61.8
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Sequence 11, Appl	Sequence 11, Appl	Sequence 57, Appl	Sequence 35, Appl	Sequence 57, Appl	•	•	•	Sequence 57, Appl	•		Sequence 14, Appl	Sequence 8, Appli	Sequence 8, Appli		Sequence 63, Appl	•	Sequence 100, App

ALIGNMENTS

US-09-025-769B-36 Patent No. 6300064 GENERAL INFORMATION: APPLICANT: Knappik APPLICANT: Pack, Pe US-09-025-769B-36 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 (EP CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/025,769B FILING DATE: 18-FEB-1998 PRIOR APPLICATION DATA: APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-FEB-1998 PRIOR APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-AUG-1995 ATTORNEY/AGENT INFORMATION: NAME: James F. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794 REFERENCE/DOCKET NUMBER: 27,794 REFERENCE/DOCKET NUMBER: MORPHO/S TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: Sequence 36, Application US/09025769B TELEFAX: (212)596-9090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 120 amino acid STRANDEDNESS: TOPOLOGY: linear MOLECULE TYPE: pro! APPLICANT: Pack, Péter APPLICANT: Ilag, Vic APPLICANT: Ge, Liming APPLICANT: Moroney, Simon APPLICANT: Moroney, Simon TITLE OF INVENTION: Protein/(Poly)peptide libraries NUMBER OF SEQUENCES: 373 CORRESPONDENCE ADDRESS: STREET: 1251 Av CITY: New York STATE: New York TYPE: amino acid COUNTRY: ADDRESSEE: 10021 New York E: James F. Haley, Jr., Esq. c/o Fish & Neave 1251 Avenue of the Americas 120 amino acids USA (212)596-9090 Knappik, Achim protein (EPO)

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VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY

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RESULT 3
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; Patent No. 6696248
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US-09-025-769B-59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/025,,
APPLICATION NUMBER: US/09/025,,
FILING DATE: 18-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95 11 3(
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., Esc
NAME: James F. Haley, Jr., MORP
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Knappi
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ge, Liming
APPLICANT: Moroney, Simon
APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                  77;
                                                                                                                                                                                                                 1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
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                                                                                                                                                                                     VQLV-QSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGWINPNSGGTNY
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                                                                                                               AQKFQGRVTMTRDTSISTAYMELSSLRSEDTAVYYCARWGGDGFYAMDYWGQGTLVTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                  120 amino acids
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Ilag, Vic
                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                   66.5%;
64.2%;
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                                                                                                                                                                                                                                                              ; Score 396.5; DB 3; ; Pred. No. 1.8e-33; 14; Mismatches 22;
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                                                                                                                                                                                                                  Sequence 59, Applicati
Patent No. 6696248
GENERAL INFORMATION:
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Best Local 8
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Versi
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/490,070A
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95 11 3021.0
PRIOR APPLICATION NUMBER: EP 95 11 3021.0
PRIOR APPLICATION NUMBER: 31.398
APPLICATION NUMBER: 31.398
ATTORNEY/AGENT INFORMATION:
NAME: Colin G. Sandercock, Esq.
REGISTRATION NUMBER: 31.398
                    Ge, Liming
Moroney, Simon
Pluckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 36:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 20006
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Colin G. Sandercock, Esq. c/o Heller Ehrman White & McAuliffe
STREET: 1666 K Street, N.W., Suite 300
CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
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Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
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                                                                                                                                                                                                                                                                                                                                                                                              61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCL-----LQALKHWGQGTLVAVSS 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 120 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ADDRESSEE: Colin G. Sandercock, Esq. c/o Heller Ehrman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (202) 912-2000
                                                                                                                                                                                                                                                               Application US/09490070A
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                                                                                                                                                   Pack, Peter
Ilag, Vic
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Ilag, Vic
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64.2%; Pred. No. 1.8e-33;
Ltive 14; Mismatches 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 36, Application US/09490153 Patent No. 6706484 GENERAL INFORMATION:
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TELEFAX: (202) 912-2020
INFORMATION FOR SEQ ID NO: 59:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/490,070A
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 77, Conserv
              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                       Moroney, Šimon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. Haley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Colin G. Sandercock, Esq. REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 37629-0005
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                         STREET: 1251 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VQLV-QSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEMMGWINPNSGGTNY 60
                                                                                                                                        COUNTRY: USA
ZIP: 10021
                                                                                                                                                                                  STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AQKFQGRVTMTRDTSISTAYMELSSLRSEDTAVYYCARWGGDGFYAMDYWGQGTLVTVSS 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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STREET: 1666 K Street, N.W., Suite 300
APPLICATION NUMBER: US/09/490,153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 20006
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                                                                                                                                                                                                                                                                                                                                                                                                     Peter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 396.5; DB 4;
Pred. No. 1.8e-33;
14; Mismatches 22;
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US-09-490-153-59
; Sequence 59, Application
; Patent No. 6706484
; GENERAL INFORMATION:
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
                                                  APPLICATION NUMBER: US/09/490,153
PILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769B
PILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
PILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                       ZIP: 10021
COMPUTER READABLE FORM:
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Ge, Liming
Moroney, Simon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
TELEPAX: (212)596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Knappik, Achim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 APKFKGRLIMSRDSSTDTVYMTLTSLTSEDTAVYYCL-----LQALKHWGQGTLVAVSS
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                                                                                                                                                                                                                                                                                                                                                                                    STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: James F. Haley, Jr., Esq. STREET: 1251 Avenue of the Americas
                  NAME: James F. Haley, Jr., REGISTRATION NUMBER: 27,79
                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: <Unknown>
REFERENCE/DOCKET NUMBER: MORPHO/5
                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: James F. Haley, Jr., REGISTRATION NUMBER: 27,794
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Pred. No. 1.8e-33;
4: Mismatches 22;
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TELECOMMUNICATION INFORMATION:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 36, Applicati Patent No. 6828422 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 59:
                                                                                                                                     INFORMATION FOR SEQ ID NO: 36:
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                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/490,324
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769
FILING DATE: 18-FEB-198
APPLICATION NUMBER: EP 95
INTERNATION NUMBER: EP 95
FILING DATE: 18-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEVERIFIED SERVICE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. Haley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                     NAME: James F. Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
TELEFAX: (212)596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plueckthun, Andreas TITLE OF INVENTION: Protein/(Poly)peptide libraries NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 36:
                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Knappik, Achim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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77; Conserv
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                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
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                                                         STRANDEDNESS: <Unknown>
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                                                                                             ENGTH: 120 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 396.5; DB 4;
Pred. No. 1.8e-33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jr., Esq.
Americas
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Version #1.30 (EPO)
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TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 59:
US-09-490-324-59
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US-09-490-324-59
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                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/490,324
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: 7,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELEPHONE: (212)596-9000
TELEPAX: (212)596-9000
INFORMATION FOR SEQ ID NO: 59:
INFORMATION FOR SEQ ID NO: 59:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 59, Application Patent No. 6828422
GENERAL INFORMATION:
                                                                                                                              Query Match
Best Local S
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                                                                                                              Matches
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Moroney, Simon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 120 amino acids
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ADDRESSEE: James F
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61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCL-----LQALKHWGQGTLVAVSS 114
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                                                          1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                              Similarity
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                                    VQLV-QSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGWINPNSGGTNY
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                                                                                                                                                                                                                                                                  TYPE: amino acid
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                                                                                                              Conservative
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Ilag, Vic
Ge, Liming
Moroney, Simon
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                                                                                                            66.5%; Score 396.5; DB 4
64.2%; Pred. No. 1.8e-33;
bive 14; Mismatches 22
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Pred. No. 1.8e-33;
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                                                                                                                                               DB 4;
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                                                                                                              Indels
                                                                                                                                             Length 120;
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RESULT 10
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Best Local
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/407
FILING DATE: 29-MAR-1995
APPLICATION NUMBER: 08/119
FILING DATE: 09-SEP-1993
APPLICATION NUMBER: 08/027
                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 05-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Olstein, Elliot M.
REGISTRATION NUMBER: 24,0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
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APPLICANT: Latinne, Dominique
TITLE OF INVENTION: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Activa
NUMBER OF SEQUENCES: 96
                                                                                                                                                                                                                                                                                                      NAME/KEY: Human Amu 5-3 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/01
FILING DATE: 07-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: New Jersey COUNTRY: U.S.A.
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                                                                                                                                 61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC-----LLQALKHWGQGTLVA 111
                                                                                                                                                                                 1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY 60
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                                            VSS
                                                                                                      AQKFQGRVTMTRDTSISTAYMELSRLRSDDTAVYYCARGRTEYIVVAEGFDYWGQGTLVT 120
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E: Cecchi, Stewart & Olstein
6 Becker Farm Road
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                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                        Score 391; DB 1; Pred. No. 7e-33;
                                                                                                                                                                                                                                                        DB 1; Length 123;
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                                                        US-08-477-989B-94
                                                                        RESULT 11
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        Sequence 94, Application US/08477989B Patent No. 5951983 GENERAL INFORMATION:
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US-08-472-281A-94
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Best Local S
Matches 75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/119,032
FILING DATE: 09-SEP-193
APPLICATION NUMBER: 08/027,008
FILING DATE: 05-MAR-193
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 08/4
FILING DATE: 29-MAR-1995
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OPERATING SYSTEM: MS-D
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
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APPLICANT: Latinne, Dominique
TITLE OF INVENTION: LO-CD2a Antibody and
                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 123 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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CITY: Roseland
'You Jers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 61'
                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Olstein, Elliot
REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER:
                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
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                                   112 VSS 114
121 VSS 123
                                                                   61 AQKFQGRVTMTRDTSISTAYMELSRLRSDDTAVYYCARGRTEYIVVAEGFDYWGQGTLVT 120
                                                                                      61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC------LLQALKHWGQGTLVA 111
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                                                                                                                                                           1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY 60
                                                                                                                                                                                                              ch 65.6%;
| Similarity 61.0%;
| 75; Conservative 1
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E: Cecchi, Stewart & Olstein
6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                                                                                                 123 amino acids
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Pred. No. 7e-33;
L6; Mismatches 22;
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Bazin, Herv

Latinne,

Dominique

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RESULT 12
US-09-025-769B-22
; Sequence 22, Application US/09025769B
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APPLICATION NUMBER: 08/407,009

FILING DATE: 29-MAR-1995

APPLICATION NUMBER: 08/119,032

APPLICATION NUMBER: 08/027,008

FILING DATE: 09-SEP-1993

APPLICATION NUMBER: 08/027,008

FILING DATE: 05-MAR-1993

ATTORNEY/AGENT INFORMATION:

NAME: 01stein, Elict M.

REGISTRATION NUMBER: 24,025
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INFORMATION FOR SEQ ID NO: 94:
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REFERENCE/DOCKET NUMBER: 61750-147
TELEPHONE 201-994-1700
TELEPHAX: 201-004
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SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,989B
FILING DATE: 07-UN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: White-Scharf, Mary
TITLE OF INVENTION: LO-CD2a Antibody and Uses
TITLE OF INVENTION: Thereof for Inhibiting
TITLE OF INVENTION: T-Cell Activation and
TITLE OF INVENTION: Proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 123 amino acids
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MEDIUM TYPE: 3.5 incl
COMPUTER: IBM PS/2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: 11:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Roseland
STATE: New Jersey
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 75; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Human Amu 5-3 heavy chain variable NAME/KEY: region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Cerella, Byrne, Bain, Gilfillan,
                                                                                                                                          112 VSS 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
                                                                                                                                                                                                                                                           VQLV-QSGAEVKKPGASVKVSCKASGYTFTGYYMHWVRQAPGQGLEWMGRINPNSGGTNY
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Kieber-Emmons, Thomas
                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                  region.
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                                                                                                                                                                                                                                                                                                                                                      Score 391; DB 2
Pred. No. 7e-33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
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APPLICANT: Knappi
APPLICANT: Pack,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212)596-9090
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: MO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., 1
REGISTRATION NUMBER: 27,794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 18-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 9:
FILING DATE: 18-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ge, Liming
APPLICANT: Moroney, Simon
APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 117 amino acids
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Pluckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
ADDRESSE: Colin G. Sandercock, Esq. c/o Helle
White & McAuliffe
                                                                                                                                                                                                                      APPLICANT: Knappik, Achim
                                                                                                                                                                                                                                                                                                                                                                        61 AQXFQGRVTMTRDTSISTAYMELSSLRSDDTAVYYCARDGDGGFDYWGQGTLVTVSS 117
                                                                                                                                                                                                                                                                                                                                                                                                         61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQ----ALKHWGQGTLVAVSS 114
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1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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Ilag, Vic
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                                                                                                                                              Pack, Peter
Ilag, Vic
Ge, Liming
Moroney, Simon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         65.3%; Score 389; DB 3; Length 117; 64.1%; Pred. No. 1.1e-32;
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                                                    Esq. c/o Heller Ehrman
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FILING DATE: 24-Jan-2000

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GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Colin G. Sandercock, Esq.
REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 37629-0005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 912-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/490,070A
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
                  CURRENT
                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Protein/(Poly)peptide libraries
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75; Conserv
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                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
                                                                                                                                                                                            ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave STREET: 1251 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AQKFQGRVTMTRDTSISTAYMELSSLRSDDTAVYYCARDGDGGFDYWGQGTLVTVSS 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VOLV-OSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGWINPNSGNTNY
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
                                                                                                                                                                          STATE: New York
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CITY: Washington
APPLICATION NUMBER: US/09/490,153
                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
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Ge, Liming
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Plueckthun, Andreas
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Pred. No. 1.1e-32;
5; Mismatches 23;
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TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: S
US-09-490-153-22
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US-09-490-324-22
; Sequence 22, A
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GENERAL INFORMATION:
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Best Local
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                                                          APPLICATION NUMBER: US/09/025,769
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                   COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                     APPLICATION NUMBER: US/09/490,324
FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
ADDRESSE: James F. Haley, Jr., Esq. c/o Fish &
STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: James F. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)596-9000
                                                                                                                                                                                                                                                                                                                                            ZIP: 10021
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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STATE: New York
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                      NAME: James F. Haley, Jr., REGISTRATION NUMBER: 27,794
                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
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REFERENCE/DOCKET NUMBER: MORPHO/5
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Ilag, Vic
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Moroney, Simon
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Pred. No. 1.1e-32;
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seq length:
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A_Geneseq_16Dec04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
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4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp20014s:*
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596
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Listing First 45 summaries
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Copyright (c) 1993 - 2005 Compugen Ltd.
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and is derived is the number of results predicted by chance to have a ster than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	ω	2	1	Result
412	413.5	413.5	414	414	418	418	418	418	419.5	421	536	536	536	536	536	564	578	588	590	596	596	596	596	596	Score
69.1	69.4	69.4	9	9	70.1		70.1	٠	•	•	•		•		89.9	•	•	98.7	•	100.0	100.0	٠	100.0		% Query Match Length
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The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and company, heart disease can be used to detect a disease as the nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides.

and coronary heart disease. Cancers include carcinogenic tumours,

Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.

Claim 30; Page 139; 163pp; English.

N-PSDB;

AAA96141.

WPI; 2000-638249/61. Ward RL, Coomber DWJ; 19-MAR-1999;

99AU-00009321.

(SVIN-)

ST VINCENT'S HOSPITAL SYDNEY LTD.

15-MAR-2000; 2000WO-AU000189

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407	407	407.5	408.5	408.5	408.5	408.5	408.5	408.5	408.5	408.5	408.5	409	409	410	410	411.5	412	412	
68.3	68.3	68.4	68.5	68.5	68.5	68.5	68.5	68.5	68.5	68.5	68.5	68.6	68.6	68.8	68.8	69.0	69.1	69.1	
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Adg96906	Abp46079	Adp03971	Adg96799	Abp45972	Adg96275			Adg96106	Abp45288		Abg76525	Adg95994	Abp45167	Adg96907	Abp46080	Adg30439	Ads92890	Ads92878	
Single ch	Human BLy	Murine-ex	Single ch	Human BLy	Single ch	Human BLy	Single ch	Single ch	Human BLy	Human BLy	HCV E1 an	Single ch	Human BLy	Single ch	Human BLy	Human GMB	Human Myo	Human Myo	

ALIGNMENTS

RESULT 1 AAB18864 p53; antibody; immune response; vaccine; gene therapy; cancer; rheumatoid arthritis; coronary heart disease. 08-FEB-2001 AAB18864 standard; protein; 28-SEP-2000. WO200056770-A1. Homo sapiens. Amino acid sequence of anti-p53 antibody heavy chain clone 163.7. AAB18864; (first entry) 114 A

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RESULT 2
AAB18860
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             The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as cuell as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectainclude carcinogenic tumours, tumours, hepatic cancer, pancreatic cancer, ovarian cancer, gastric cancer, brain cancer, brain cancer, brain cancer, prostate cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 114;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                 Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
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N-PSDB; AAA96137.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-MAR-1999;
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   cancer,
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   oesophageal cancer,
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Pred. No. 1.1e-46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ine; gene therapy; cancer, disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 114;
 mesenchymal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60
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RESULT 3
AAB18876
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                                     The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in correctnes, for inducing an immune response against a disease in a correctnes, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as computed as the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer, tumours, the disease cancer, pancreatic cancer, brain cancer, bladder cancer, prostate cancer and cancer, brain cancer, bladder cancer, mesenchymal tumours, and heart and heart cancer, brain cancer, pancreatic cancer, mesenchymal tumours,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  p53; antibody; immune
rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                           e.g. sarcoma,
                                                                                                                                                                                                                                                                                                                Claim 30; Page 151; 163pp; English
                                                                                                                                                                                                                                                                                                                                                            Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ward RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200056770-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coomber DWJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
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                             and
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                             hemopoietic tumours, e.g. B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of anti-p53 antibody heavy chain clone 163.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  response; vaccine; gene therapy; cancer; coronary heart disease.
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                                                                                                                                                                                                                                                                                                                                                   disease
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Pred. No. 1.1e-46;
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                           cell
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                             Lymphoma
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AAB18862
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         Query Match
Best Local Sim
Matches 114;
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Best Local
                                                                                                                                                                                                                                                                          The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, head and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer,
                                                                                                                                                                                    head and neck tumours, hepatic cancer, pancreatic cancer, ovarian gastric cancer, brain cancer, bladder cancer, prostate cancer and urinary/genital tract cancer, oseophageal cancer, mesenchymal tumes.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 30; Page 137; 163pp; English
                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatoid arthritis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arthritis and coronary heart disease.
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                                         Local Similarity
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Score 596; Db 3;
Pred. No. 1.1e-46;
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Pred. No. 1.1e-46;
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문 5

Matches

Similarity

Conservative

VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY

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RESULT 5
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                                                                                                  The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain C chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a corresponse. The nucleic acid sequences can be used to detect and for detection purposes. The nucleic acid sequences can be used to detect a disease as computed to the polypeptides. In particular, the following can be treated cancer, rhematoid arthritis and coronary heart disease. Cancer include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, chead and neck tumours, hepatic cancer, postate cancer overian cancer, castric cancer, brain cancer, bladder cancer, prostate cancer and curinary/genital tract cancer; oseophageal cancer, mesenchymal tumours, ce.g. e.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma
                    Query Match
Best Local
                                                                          Sequence 114
                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 30; Page 141; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
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rheumatoid arthritis;
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100.0%; E
100.0%; E
tive 0;
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coronary heart
Score 596; I
Pred. No. 1.1
); Mismatches
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heart disease.
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= 596; DB 3;
. No. 1.1e-46;
smatches 0;
                                 Length 114;
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61

APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS 114

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RESULT 6
AAB18868
ID AAB1
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                                                                                                                                        Matches
                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, head and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, gastric cancer, brain cancer, bladder cancer, prostate cancer and urinary/genital tract cancer, oesophageal cancer, mesenchymal tumours, e.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma
                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelacors, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     p53; antibody; immune rheumatoid arthritis;
                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ward RL,
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                                                                                                                                        113;
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 61
                                  61
                                                                    \vdash
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                              APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS 114
                                                                                         VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                                                                                                        114 AA;
APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 143; 163pp;
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                                                                                                                                        Conservative
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                                                                                                                                                      99.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       response; vaccine; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
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                                                                                                                                      Score 590; DB 3
Pred. No. 4e-46;
0; Mismatches
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                                                                                                                                                                      Length 114;
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                                                                                                                                                                                                                                                                                                                               tumours
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RESULT 8 AAB18874 ID AAB1

AAB18874 standard; protein; 114

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RESULT 7
AAB18858
ID AAB1
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                                                                                                                                   Query Match
Best Local Similarity
Matches 112; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polynucleotides encoding anti-\mathfrak{p}53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
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                                                                                                                                                                                                        Sequence 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          p53; antibody; immune response; vaccine; gene therapy; cancer;
rheumatoid arthritis; coronary heart disease.
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DB; AAA96135.
19
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                                                                                                 2 QLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYA
                                  PKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS 114
                                                                  QLLEQSGAEVKRPGASVTISCRASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANYA
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                                                                                                                                     Conservative
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                                                                                                                                                     98.7%;
99.1%;
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                                                                                                                                   Score 588; DB 3;
Pred. No. 6.1e-46;
1; Mismatches 0
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                                                                                                                                                                   Length 114;
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 113
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                                                                                                                                        Best Loc
Matches
                                                                                                                                                                         Query Match
                                                                                                                                                                                                            Sequence
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rheumatoid arthritis; coronary heart disease.
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                                                                                                                                                          Local
                                                                                                                                        al Similarity
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                                 51
                                                                    \vdash
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                    APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS
                                                                VQLLEQSGAEVKRPGASVTISCRASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                        VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
APKFKGRLTMSRDSSTETVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVGPSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence
                                                                                                                                          Conservative
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Fi
                                                                                                                                                       97.0%;
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                                                                                                                                      Score 578; DB 3
Pred. No. 5e-45;
2; Mismatches
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RESULT 9 AAB18882 ID AAB1

AAB18882 standard; protein; 114 AA

08-FEB-2001 AAB18880;

(first entry)

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RESULT 10
AAB18880
ID AAB18
XX
AC AAB18
XX
DT 08-FE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 106;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                             AAB18880 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 114 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polynucleotides encoding anti-p53 antibodies, polypeptides and pefragments, useful in treatment and diagnosis of cancer, rheumatoi arthritis and coronary heart disease.
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rheumatoid arthritis; coronary heart disease.
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                                                                                                                                                                                                                                                                                                                                                             VQLLEQSGAEVKRPGASVTISCQASRQNFSGQYIHWVRQAPGQGLEWMGIINPSGGSANY
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                                                                                                          protein; 114
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93.0%;
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Pred. No. 9.5e-44;
6; Mismatches 2
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
p53; antibody; immune response; vaccine; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 114 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain
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N-PSDB; AAA96157.
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                                                                 Amino acid sequence of anti-p53 antibody heavy chain clone 163.22.
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                                                                                                                                                                                                                                                           standard;
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Pred. No. 3.4e-41;
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RESULT 12
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Best Local (
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                                                                                           p53; antibody; immune response; vaccine; gene
rheumatoid arthritis; coronary heart disease.
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Pred. No. 3.4e
8; Mïsmatches
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No. 3.4e-41;
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                28-SEP-2000
                                                                                                                 p53; antibody; immune response; vaccine; gene therapy; cancer;
rheumatoid arthritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 30; Page 145; 163pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-638249/61.
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                                                  WO200056770-A1
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                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        101;
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                                                                                                                                                                                                                                                                                                                                                                  13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                                                                                                                                                                                                                                                                                                                    APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQALKHWGQGTLVAVSS 114
                                                                                                                                                                                                                                                                                                                                                                APSFQGRLSMSRDASTNTVYMKLSSLTSEDTAVYYCLSQALKYWGQGTLVAVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                   VQLLEQSGAEMKRPGASVTISCQASRQTFSGQYIHWVRQAPGQGLEWMGVINPSGGSANY
                                                                                                                                                                      sequence
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                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                      of anti-p53 antibody heavy chain clone 163.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 536; DB 3; pred. No. 3.4e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>ა</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 114;
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RESULT 14
AAB11872
ID AAB18
XX AAB18
AC AAB18
XX AAB18
DT 08-FE
XX AMING
XX P53;
KW P53;
KW P53;
KW P53;
KW P53;
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KW P64m
XX P6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          an immune response against a naturally occurring disease. The antibodies chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a coronary treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as cell as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, the patic cancer, pancreatic cancer, ovarian cancer, cancer gastric cancer, brain cancer, bladder cancer, prostate cancer and curinary/genital tract cancer, bladder cancer, mesenchymal tumours, and heave tumours and heave tumours and heave tumours, tempoure and and neck tumours head and neck tumours, hepatic cancer, prostate cancer and cancer, pastric cancer and tract cancer, bladder cancer, mesenchymal tumours, and heave tumours and heave tumours and heave the cancer and cancer, cancer and cancer and cancer, cancer and can
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Best Local
                                              19-MAR-1999;
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                                                                                                                     15-MAR-2000; 2000WO-AU000189
                                                                                                                                                                                                         28-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 p53; antibody; immune response; vaccine; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amino acid sequence of anti-p53 antibody heavy chain clone 163.16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB18872;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 114 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 30; Page 131; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               rheumatoid arthritis; coronary heart disease.
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DB; AAA96133.
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Pred. No. 3.4e-41;
8; Mismatches 5
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents the heavy chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease.
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                                                                                                                                             Mus sp.
Synthetic.
                                                                                                                                                                                                        Anti-murine CTLA-4 sFv; M19 sFv; single chain antibody; murine CTLA4; membrane-associated protein; chimeric construct; extracellular domain; human CD8; ligand; activated T-cell; co-stimulatory signal; donor B7; recipient CD28; T-cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in
                 11-NOV-1999
                                                                                                                                                               Mug
                                                                                                                                                                                            xenograft-specific
                                                                                                                                                                                                                                                                                             Anti-murine CTLA-4 M19 sFv.
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                                               WO9957266-A2
                                                                                               Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                           standard;
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                                                                                             Location/Qualifiers 208
                                                                               note=
                                                                                                                                                                                                                                                                                                                                                                                         protein; 242
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                                                                                                                                                                                           immunosuppression.
                                                                              "Corresponds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
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                                                                              atc codon"
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Best Local S
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                                                                                                                                                                                                              Sequence 242
                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                             Inhibiting T-cell mediated rejection
                                                                                                                                                                                                                                         xenograft-specific immunosuppression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lechler IR,
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                                         61
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                                                                                                                 1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
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                                                                                                                                                                     Similarity
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                                         AQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARVAPYVNTLVFWGQGTLVTVSS
                                                      APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC-----LLQALKHWGQGTLVAVSS
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2005, 05:59:09
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12; Mismatches
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Search completed: November Job time : 168 secs

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Maximum DB seq length: 200000000
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Published Applications AA:*

1: /cgn2_6/ptcdata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptcdata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptcdata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptcdata/1/pubpaa/US06_PUBCOMB.pep:*

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6: /cgn2_6/ptcdata/1/pubpaa/US08_NEW_PUB.pep:*

7: /cgn2_6/ptcdata/1/pubpaa/US08_NEW_PUB.pep:*

8: /cgn2_6/ptcdata/1/pubpaa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptcdata/1/pubpaa/US09B_PUBCOMB.pep:*

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12: /cgn2_6/ptcdata/1/pubpaa/US09C_PUBCOMB.pep:*

13: /cgn2_6/ptcdata/1/pubpaa/US10A_PUBCOMB.pep:*

14: /cgn2_6/ptcdata/1/pubpaa/US10A_PUBCOMB.pep:*

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17: /cgn2_6/ptcdata/1/pubpaa/US10A_PUBCOMB.pep:*

18: /cgn2_6/ptcdata/1/pubpaa/US10C_PUBCOMB.pep:*

19: /cgn2_6/ptcdata/1/pubpaa/US10R_PUBCOMB.pep:*

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25: /cgn2_6/ptcdata/1/pubpaa/US10R_PUB.pep:*
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596
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Regult No.	Score	Query Match	Query Match Length DB	DB	ID	Description
1	418	70.1	142	14	US-10-171-452A-2	Sequence 2, Appli
ນ	418	70.1	142	15	US-10-353-708-2	Sequence 2, Appli
ω	418	70.1	142	16	US-10-731-984-36	Sequence 36, Appl
4	414	69.5	255	10	US-09-880-748-1156	Sequence 1156, Ap
v	414	69.5	255	15	US-10-293-418-1156	Sequence 1156, Ap
σ	413.5	69.4	245	10	US-09-880-748-1919	Sequence 1919, Ap
7	413.5	69.4	245	15	US-10-293-418-1919	
89	412	69.1	117	16	US-10-688-925-16	
9	412	69.1	117	16	US-10-688-925-28	28
10	412	69.1	249	16	US-10-688-925-14	14
11	412	69.1	249	16	US-10-688-925-26	Sequence 26, Appl

ALIGNMENTS

TYPE: PRI TYPE: PRI ORGANISM: Homo sapiens -10-171-452A-2	ID NO 2	OF SEC ID NOS: 60	APPLICATION N	FILING DATE: 2001-09-20	FILING DATE: 2002-10-19 APPLICATION NUMBER: GB012272	APPLICATION N	FILING DATE: 2002-04	FILING DATE: 2002-04	APPLICATION NUMB	FILING DATE: 2003-0	NT APPLICATION NUMBER:	RENCE: 695458-59	THUENTION: TBY1 Antibody and Hoos	ш	: Hale, Ge	: Gorman, Sc	Waldmann,	'n	GENERAL INFORMATION:	Publication No. US20030108518A1	Sequence 2, Application US/10171452A	-10-171-452A-2	מטכני ד
WINSOR-HINES, DAWN NVENTION: TRX1 ANTIDODY and Uses ENCE: 695458-59 FELICATION NUMBER: US/10/171,452A LING DATE: 2003-02-10 LCATION NUMBER: US60/373,471 NG DATE: 2002-04-18 LCATION NUMBER: US60/373,470 NG DATE: 2002-04-18 LCATION NUMBER: US60/345,194 NG DATE: 2002-10-19 LCATION NUMBER: US60/345,194 NG DATE: 2002-10-19 LCATION NUMBER: GB0122724.8 NG DATE: 2001-09-20 LCATION NUMBER: GB0127274.8 NG DATE: 2001-09-20	CAVI: WINSOT-HINES, DAWN OF INVENTION: TRXI ARTHODY and USES REFERENCE: 695458-59 WIT APPLICATION NUMBER: US/10/171,452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0122724.8 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0127724.8 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6	CAVI: WINSOT-HINES, DAWN OF INVENTION: TRXI ARTIDODY and USES REFERENCE: 695458-59 NUT APPLICATION NUMBER: US/10/171,452A NUT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: GB0122724-8 APPLICATION NUMBER: GB0122724-8 APPLICATION NUMBER: GB0124724-8 APPLICATION NUMBER: GB0114517-6	CAVI: WINSOT-HINES, DAWN OF INVENTION: TRXI ARTIDODY and Uses REFERENCE: 695458-59 WIT APPLICATION NUMBER: US/10/171,452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 APPLICATION NUMBER: US60/345,194 FILING DATE: 2001-10-19 APPLICATION NUMBER: US60/345,194	CAN': WINSOT-HINES, DAWN OF INVENTION: TRXI ARTIDODY and Uses REFERENCE: 695458-59 NIT APPLICATION NUMBER: US/10/171,452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: GROIN-12774-8 APPLICATION NUMBER: GROIN-12774-8	CAVI: WINSOT-HINES, DAWN OF INVENTION: TRXI ANTIDODY and USES REFERENCE: 695458-59 NIT APPLICATION NUMBER: US/10/171,452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US/60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US/60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US/60/345,194	INVENTION: TRX1 Antibody and Uses RENCE: 695458-59 PPLICATION NUMBER: US/10/171,452A TLING DATE: 2003-02-10 LICATION NUMBER: US60/373,471 LICATION NUMBER: US60/373,470	INVENTION: TRX1 Antibody and Uses RENCE: 695458-59 TEPLICATION NUMBER: US/10/171,452A TLING DATE: 2003-02-10 LICATION NUMBER: US60/373,471 LICATION SUMBER: US60/373,471 LICATION SUMBER: US60/373,471	INVENTION: TRX1 Antibody and Uses RENCE: 695458-59 IPPLICATION NUMBER: US/10/171,452A TLING DATE: 2003-02-10 TICATION NUMBER: US60/373,471	. WINSOY-HINES, DAWN INVENTION: TRX1 Antibody and Uses RENCE: 695458-59 PPLICATION NUMBER: US/10/171,452A ILING DATE: 2003-02-10	:: winsor-Hines, Dawn INVENTION: TRX1 Antibody and RENCE: 695458-59 PPLICATION NUMBER: US/10/171.	RENCE: 695458-59	THUENTION: TBY1 Antibody and Hoos		Kornaga, Ringler,	Rao, Patr Kornaga, Ringler,	: Hale, Geo : Rao, Patr : Kornaga, : Ringler,	: Gorman, S : Hale, Geo : Rao, Patr : Kornaga, : Ringler,	: Waldmann, : Gorman, S : Hale, Geo : Rao, Patr : Kornaga, : Ringler,	: Frewin, Ma : Waldmann, S : Gorman, S : Hale, Geo : Rao, Patr : Kornaga, : Ringler,	GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricía APPLICANT: Kornaga, Tadeusz APPLICANT: Ringler, Douglas	Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Rro, Patricia APPLICANT: Kornaga, Tadeusz APPLICANT: Ringler, Douglas	plicatio . US2003 ATION: ewin, Ma aldmann, orman, S ale, Geo ao, Patr ornaga, ingler,	S-10-171-452A-2 Sequence 2, Application US/10171452A Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Waldmann, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Kornaga, Tadeus APPLICANT: Kornaga, Tadeus APPLICANT: Ringler, Douglas
PLICANT: Cobbold, Stephen PLE ANT: Winsor-Hines, Dawn PLE OF INVENTION: TEX1 Antibody and Uses REFERENCE: 695458-59 REENT APPLICATION NUMBER: US/10/171,452A RENT APPLICATION NUMBER: US60/373,471 FOR FILING DATE: 2002-04-18 FOR APPLICATION NUMBER: US60/373,470 FOR FILING DATE: 2002-04-18 FOR FILING DATE: 2002-04-18 FOR APPLICATION NUMBER: US60/345,194 FOR FILING DATE: 2002-10-19 FOR FILING DATE: 2001-09-20 FOR FILING DATE: 2001-06-14 FILING DATE: 2001-06-14 FOR FILING DATE: 60 FOR F	CANT: Cobbold, Stephen CANT: Winsor-Hines, Dawn OF INVENTION: TRX1 Antibody and Uses OF INVENTION: TRX1 Antibody and Uses NT REFERENCE: 695458-59 NT APPLICATION NUMBER: US/10/171,452A NT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: GB0127724.8 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6	CANT: Cobbold, Stephen CANT: Winsor-Hines, Dawn OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NT APPLICATION NUMBER: US/10/171, 452A NT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: GB0127724-8 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB01127721-6 APPLICATION NUMBER: GB01127721-6	CANT: Cobbold, Stephen CANT: Winsor-Hines, Dawn OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NIT APPLICATION NUMBER: US/10/171, 452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-09-20 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Kornaga,</pre>	: Gorman, S : Hale, Geo : Rao, Patr : Kornaga,	: Waldmann, : Gorman, S : Hale, Geo : Rao, Patr : Kornaga,	: Frewin, Ma : Waldmann, : Gorman, S : Hale, Geo : Rao, Patr : Kornaga,	GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Kornaga, Tadeusz	Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Roo, Patricia	Sequence 2, Application US/10171452A Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Roormaga, Tadeusz	S-10-171-452A-2 Sequence 2, Application US/10171452A Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia APPLICANT: Rao, Patricia
PLICANT: Ringler, Douglas PLICANT: Winsor-Hines, Dawn PLICANT: Winsor-Hines, Dawn PLE OF INVENTION: TRX1 Antibody and Uses LE REFERENCE: 695458-59 REENT APPLICATION NUMBER: US/10/171,452A REENT APPLICATION NUMBER: US60/373,471 FOR APPLICATION NUMBER: US60/373,471 FOR APPLICATION NUMBER: US60/373,470 FOR FILING DATE: 2002-04-18 FOR FILING DATE: 2002-04-18 FOR FILING DATE: 2002-10-19 FOR FILING DATE: 2002-10-19 FOR FILING DATE: 2002-10-19 FOR FILING DATE: 2001-09-20 FOR APPLICATION NUMBER: GB012774.8 FOR FILING DATE: 2001-09-20 FOR APPLICATION NUMBER: GB012774.8 FOR FILING DATE: 2001-09-20 FOR APPLICATION NUMBER: GB012774.8 FOR APPLICATION NUMBER: GB012774.8 FOR FILING DATE: 2001-09-20 FOR APPLICATION NUMBER: GB012774.8 FOR APPLICATION	CANT: Ringler, Douglas CANT: Cobbold, Stephen OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NT APPLICATION NUMBER: US/10/171,452A NT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: US60/345,194 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6	CANT: Ringler, Douglas CANT: Cobbold, Stephen OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NIT APPLICATION NUMBER: US/10/171,452A NIT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-8 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: US60/345,194 FILING DATE: 2002-10-19 APPLICATION NUMBER: US60/345,194 FILING DATE: 2001-09-20 APPLICATION NUMBER: GB012774.8 BFILING DATE: 2001-09-20 APPLICATION NUMBER: GB0114517.6	CANT: Ringler, Douglas CANT: Cobbold, Stephen CANT: Cobbold, Stephen OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NT APPLICATION NUMBER: US/10/171, 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GR0127774-8	CANT: Ringler, Douglas CANT: Cobbold, Stephen CANT: Cobbold, Stephen CANT: Winsor-Hines, Dawn OF INVENTION: TRX1 Antibody and Uses REFERENCE: 695458-59 NT APPLICATION NUMBER: US/10/171,452A NT FILING DATE: 2003-02-10 APPLICATION NUMBER: US60/373,471 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 APPLICATION NUMBER: US60/373,470 FILING DATE: 2002-04-18 APPLICATION NUMBER: US60/373,470 APPLICATION NUMBER: US60/373,470	: Ringler, Douglas : Cobbold, Stephen : Winsor-Hines, Dawn INVENTION: TRX1 Antibody and Uses PPLICATION NUMBER: US/10/171,452A 'LING DATE: 2003-02-10 LICATION NUMBER: US60/373,471 LIG DATE: 2002-04-18 LICATION NUMBER: US60/373,470	: Ringler, Douglas : Cobbold, Stephen : Winsor-Hines, Dawn INVENTION: TRX1 Antibody and Uses PPLICATION NUMBER: US/10/171,452A TLING DATE: 2003-02-10 LING DATE: 2002-04-18	: Ringler, Douglas : Cobbold, Stephen : Winsor-Hines, Dawn INVENTION: TRX1 Antibody and Uses IRENCE: 695458-59 PPLICATION NUMBER: US/10/171,452A TILING DATE: 2003-02-10 LICATION NUMBER: US60/373,471	: Ringler, Douglas : Cobbold, Stephen : Winsor-Hines, Dawn INVENTION: TRX1 Antibody and Uses RENCE: 69458-59 PPLICATION NUMBER: US/10/171,452A TLING DATE: 2003-02-10	: Ringler, Douglas : Cobbold, Stephen : Winsor-Hines, Dawn : Winsor-Hines, Wins	Ringler, Douglas Cobbold, Stephen Minsor-Hines, Dawn INVENTION: TRX1 Antibody and Uses RENCE: 695458-59	Ringler, Douglas Cobbold, Stephen Dawn TANDANTON TOTAL	Ringler, Dougl Cobbold, Steph		: Rao,	: Hale, Ge : Rao, Pat	<pre>: Gorman, Sc : Hale, Geof : Rao, Patri</pre>	: Waldmann,: Gorman, Sc: Hale, Geof: Rao, Patri	<pre>: Frewin, Max : Waldmann, : Gorman, Sc : Hale, Geof : Rao, Patri</pre>	GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Hale, Geoff	Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Hac, Geoff	Sequence 2, Application US/10171452A Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Hale, Geoff APPLICANT: Rao, Patricia	S-10-171-452A-2 Sequence 2, Application US/10171452A Publication No. US20030108518A1 GENERAL INFORMATION: APPLICANT: Frewin, Mark APPLICANT: Waldmann, Herman APPLICANT: Gorman, Scott APPLICANT: Hale, Geoff APPLICANT: Hale, Geoff

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; NUMBER OF SEQ ID NOS: 60
; SEQ ID NO 2
; LENGTH: 142
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-353-708-2
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                                                                                                                                                                                                                                                      Query Match 70.1%; Score 418; DB 15; Best Local Similarity 63.8%; Pred. No. 2.7e-31; Matches 81; Conservative 15; Mismatches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/10353708 Publication No. US20030219403A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Winsor-Hines, Dawn
TITLE OF INVENTION: Compositions and Methods of Tolerizing a Primate to
FILE REFERENCE: 695458-73
CURRENT APPLICATION NUMBER: US/10/353,708
CURRENT FILING DATE: 2003-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US60/373,470 PRIOR FILING DATE: 2002-04-18 PRIOR APPLICATION NUMBER: US60/345,194 PRIOR FILING DATE: 2002-10-19 PRIOR APPLICATION NUMBER: GB0122724.8 PRIOR FILING DATE: 2001-09-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Frewin, Mark
APPLICANT: Waldmann, He
APPLICANT: Gorman, Scot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US10/171,452
PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US60/373,471
PRIOR FILING DATE: 2002-04-18
PRIOR FILING DATE: 2002-04-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: GB0114517.6 PRIOR FILING DATE: 2001-06-14
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                                       108 TLVAVSS 114
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                                                                                    71
                                                                                                        61 APKFKGRLIMSRDSSTDTVYMTLTSLTSEDTAVYYC------LQALKHWGQG
                                                                                                                                                                 12 VQLV-QSGAEVKKPGASVKVSCKASGYTFTNYYMHWVRQAPGQGLEWMGIINPSGNSTNY 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYC------LLQALKHWGQG 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  81; Conservative
                                                                                                                                                                                              1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
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TLVTVSS 137
                                                                                  AQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCAREKLATTIFGVLIITGMDYWGQG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VQLV-QSGAEVKKPGASVKVSCKASGYTFTNYYMHWVRQAPGQGLEWMGIINPSGNSTNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ringler, Douglas
Cobbold, Stephen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rao, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Waldmann, Her
Gorman, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Geoff
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        70.1%; Score 418; DB 14; 63.8%; Pred. No. 2.7e-31;
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                                                                                                                          107
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US-09-880-748-1156
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                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
PRIOR PILING DATE: 2000-06-15
PRIOR PLING DATE: 2000-10-17
PRIOR PLING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-21
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
SOFTWARE: PATENTIN VET: 2.0
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                                                                                              US-09-880-748-1156
                                                                                                                                   SEQ ID NO 1156
LENGTH: 255
TYPE: PRT
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Matches
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  Matches
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                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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APPLICANT: RINGLER, Douglas J.
APPLICANT: RINGLER, Douglas J.
APPLICANT: RINGLER, DOUGLING TOLERANCE IN PRIMATES
FILE REFERENCE: TLN-022
CURRENT APPLICATION NUMBER: US/10/731,984
CURRENT FILING DATE: 2003-12-09
PRIOR APPLICATION NUMBER: 60/431839
PRIOR FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 70
NUMBER OF SEQ ID NOS: 70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: WINDSOR-HINES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: PF523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Synthetic Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81;
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  Conservative
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                        69.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   70.1%; Score 418; DB 16; 63.8%; Pred. No. 2.7e-31;
Score 414; DB 10;
Pred. No. 1.2e-30;
4; Mismatches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                           Length 255;
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  Indels 14;
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  Gaps
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RESULT 3

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US-09-880-748-1919
; Sequence 1919, Application US/09880748
; Publication No. US20030059937A1
; GENERAL INFORMATION:
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ORGANISM: Homo sapiens
US-10-293-418-1156
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                                                                                                                                                                                                  RESULT 6
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PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/297,379
PRIOR APPLICATION NUMBER: 60/277,379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 1156
LENGTH: 255
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Best Local !
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GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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PRIOR APPLICATION NUMBER: 60/2
PRIOR FILING DATE: 700ER: 60/2
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 69.5%; Score 414; DB 15;
Similarity 63.2%; Pred. No. 1.2e-30;
79; Conservative 14; Mismatches 18;
                                                                                                                                                                                                                                                                                                                                                                       FKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCL-----LQALKHWGQGTL 109
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                                                                                                                                                                                                                                                                                                                                   FQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARGGEYDILTGYYFGLGVYDYWGQGTM 123
                                                                                                                                                                                                                                                                                                                                                                                                             LQQSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGIINPSGGSTSYAQK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FOGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARGGEYDILTGYYFGLGVYDYWGQGTM 123
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. US20030223996A1
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                                                                                                                                        US-10-293-418-1919
                                                            Query Match
Best Local S
Matches 81
                                                                                                                                                                                                                  SEQ ID NO 1919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PF523P2
                                                                                                                                                                              LENGTH: 245
TYPE: PRT
                                                                                                                                                          ORGANISM: Homo sapiens
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1 VQLLEQSGAEVKRPGASVTISCQASRQDFSGQYIHWVRQAPGQGFEWMGIINPSGGSANY
                                                                                Similarity
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; LENGTH: 245
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-880-748-1919
                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR PPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 3247
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Publication No. US20030223996A1
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Best Local Similarity 66.4%; Pred. No. 1.2e-30;
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PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR FILING DATE: 2001-06-15
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CURRENT FILING DATE: 2002-11-27
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PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
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NUMBER OF SEQ ID NOS: 3239
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PRIOR FILING DATE: 2001-05-25
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FILING DATE: 2001-05-25
APPLICATION NUMBER: 60/277,379
FILING DATE: 2001-03-21
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       Conservative
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69.4%; Score 413.5; DB 15 66.4%; Pred. No. 1.2e-30; tive 14; Mismatches 18;
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                                                     DB 15;
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Sequence 28, Application US/10688925
Publication No. US20040142382A1
GENERAL INFORMATION:
APPLICANT: Veldman, Geertruida et al.
TITLE OF INVENTION: NEUTRALIZING ANTIBODIES AGAINST GDE
FILE REFERENCE: 08702.0020-00000
CURRENT APPLICATION NUMBER: US/10/688,925
CURRENT FILING DATE: 2003-10-21
NUMBER OF SEQ ID NOS: 54
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 28
                                                                                                                                                                                                  ; LENGTH: 117
; TYPE: PRT
; ORGANISM: Homo &
US-10-688-925-28
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Sequence 16, Application US/10688925
Publication No. US20040142382A1
GENERAL INFORMATION:
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SEQ ID NO 16

LENGTH: 117

TYPE: PRT

ORGANISM: Homo sapiens

US-10-688-925-16
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Best Local Similarity
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TITLE OF INVENTION: NEUTRALIZING ANTIBODIES
FILE REFERENCE: 08702.0020-000000
CURRENT APPLICATION NUMBER: US/10/688,925
CURRENT FILING DATE: 2003-10-21
NUMBER OF SEQ ID NOS: 54
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                               APKFKGRLTMSRDSSTDTVYMTLTSLTSEDTAVYYCLLQ---ALKHWGQGTLVAVSS 114
AQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARDENWGFDPWGQGTLVTVSS 117
                                                               VQLV-QSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGIINPSGGSTSY
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Pred. No. 8e-31;
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Pred. No. 8e-31;
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US-10-935-290-72
Sequence 72, Application US/10935290
Publication No. US20050069542A1
GENERAL INFORMATION:
APPLICANT: Baker et al.
TITLE OF INVENTION: Antibodies that Specifically Bind to GMAD
FILE REFERENCE: PF584P1
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SEQ ID NO 26
; LENGTH: 249
; TYPE: PRT
; ORGANISM: Homo sapiens
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US-10-688-925-26
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Publication No. US20040142382A1

GENERAL INFORMATION:
APPLICANT: Voldman, Geextruida et al.
TITLE OF INVENTION: NEUTRALIZING ANTIBODIES AGAINST GDF
FILE REFERENCE: 08702.0020-00000

CURRENT APPLICATION NUMBER: US/10/688,925

CURRENT FILING DATE: 2003-10-21

NUMBER OF SEQ ID NOS: 54

SOFTWARE: Patentin version 3.1
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LENGTH: 249
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Best Local
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TITLE OF INVENTION: NEUTRALIZING ANTIBODIES AGAINST GDF 8
FILE REFERENCE: 08702.0020-00000
CURRENT APPLICATION NUMBER: US/10/688,925
CURRENT FILING DATE: 2003-10-21
NUMBER OF SEQ ID NOS: 54
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CURRENT APPLICATION NUMBER: US/10/935,290 CURRENT FILING DATE: 2004-09-08 PRIOR APPLICATION NUMBER: PCT/US03/09625
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Pred. No. 1.7e-30;
2; Mismatches 20
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Pred. No. 1.7e-30;
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RESULT 14
US-10-293-418-2091
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SEQ ID NO 2091
LENGTH: 248
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PRIOR FILING DATE: 2000-06-15
PRIOR PELICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
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PRIOR APPLICATION NUMBER: 60/368,813
PRIOR FILING DATE: 2002-04-01
NUMBER OF SEQ ID NOS: 234
SEQ ID NO 72
LENGTH: 241
TYPE: PRT
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CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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1 Similarity 66.9%;
79; Conservation
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   Application US/10293418
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                                                                                                                                                                                                                                                                                                                      68.8%; Score 410; DB 10; 65.3%; Pred. No. 2.7e-30;
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CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/21,210
PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR PPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR PILING DATE: 2001-05-25
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PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/280,748
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
                                                                       SOFTWARE: PatentIn Ver. SEQ ID NO 1178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Ruben et al
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Best Local
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SEQ ID NO 2091
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
                                                                                                                        NUMBER OF SEQ ID NOS: 3239
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TYPE: PRT
ORGANISM: Homo sapiens
ORGANISM: Homo sapiens
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FILING DATE: 2001-03-21
APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
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Pred. No. 2.7e-30;
7; Mismatches 17;
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